RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: ___*10/612,468*___
Source: ___*OIPE*___
Date Processed by STIC: ___*7/29/2003*___

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PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**
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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , EFS Submission **User Manual** - ePAVE)
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Revised 04/24/2003

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OIPE

RAW SEQUENCE LISTING   DATE: 07/29/2003
PATENT APPLICATION: US/10/612,468   TIME: 13:35:30

Input Set : A:\EP.txt
Output Set: N:\CRF4\07292003\J612468.raw

```
        3 <110> APPLICANT: Zhang, Jingwu Z.
        4        Ho, Walter Kowk Keung
        5        Zhang, Dongqing
        6        Sun, Wei
W-->    7 <120> TITLE OF INVENTION: T Cell Receptor CDR3 Sequence and Methods for Detecting and
Treating
W-->    8        Rheumatoid Arthritis
W-->    9 <130> FILE REFERENCE: 057186.000003
W-->   10 <140> CURRENT APPLICATION NUMBER:
C-->   11 <141> CURRENT FILING DATE: 2003-07-02      >< 1507  ← insert this mandatory numeric
W-->   11 <151> PRIOR FILING DATE: 2002-07-02                    identifier and its response
W-->   12 <160> NUMBER OF SEQ ID: 168                                 wherever < 151>

ERRORED SEQUENCES                                         Does Not Comply       for a
                                                          Corrected Diskette Needed   response
      775 <210> SEQ ID NO: 66                             pp 1,3
      776 <211> LENGTH: (21) 19 shown below
      777 <212> TYPE: DNA
      778 <213> ORGANISM: Artificial
W-->  779 <220> FEATURE:
      780 <221> NAME/KEY: primer_bind
      781 <223> OTHER INFORMATION: FAM (expand)-labeled BJ primer used for run-off reactions
W-->  782 <400> SEQUENCE: 66
E-->  784 ggagagtcga gttccatca                             19
     1416 <210> SEQ ID NO: 114
     1417 <211> LENGTH: 63
     1418 <212> TYPE: DNA
     1419 <213> ORGANISM: Artificial
W-->  1420 <220> FEATURE:
W-->  1421 <221> NAME/KEY:
     1422 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
     1423        RA patients
GK->  1424 <400> SEQUENCE: 114
     1426 tacttctgtg ccagcagtcc cacgcgggac aggggaaata atgagcagtt cttcgggcca    60
E-->  1427 gga                                                          63 ← insert
     1445 <210> SEQ ID NO: 116
     1446 <211> LENGTH: 63
     1447 <212> TYPE: DNA
     1448 <213> ORGANISM: Artificial    p.3
W-->  1449 <220> FEATURE:
W-->  1450 <221> NAME/KEY:
     1451 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
```

ST specimen of
    1452   RA patients

7/29/03

W--> 1453 <400> SEQUENCE: 116
     1455 tacttctgtg ccagcagttc cccaatagcg gggagctcca atgagcagtt cttcgggcca   60
E--> 1456 gga   63 ← insert
     1473 <210> SEQ ID NO: 118
     1474 <211> LENGTH: 63
     1475 <212> TYPE: DNA
     1476 <213> ORGANISM: Artificial
W--> 1477 <220> FEATURE:
W--> 1478 <221> NAME/KEY:
     1479 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
     1480   RA patients
W--> 1481 <400> SEQUENCE: 118
     1483 tacttctgtg ccagcagttt ctgggcccct acggacaata atgagcagtt cttcgggcca   60
E--> 1484 gga   63 ← insert
     1529 <210> SEQ ID NO: 122
     1530 <211> LENGTH: 63
     1531 <212> TYPE: DNA
     1532 <213> ORGANISM: Artificial
W--> 1533 <220> FEATURE:
W--> 1534 <221> NAME/KEY:
     1535 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
     1536   RA patients
W--> 1537 <400> SEQUENCE: 122
     1539 tacttctgtg ccagcagccc tagggagggc ctcctcaata atgagcagtt cttcgggcca   60
E--> 1540 gga   63 ← insert
     1888 <210> SEQ ID NO: 148
     1889 <211> LENGTH: 61
     1890 <212> TYPE: DNA
     1891 <213> ORGANISM: Artificial
W--> 1892 <220> FEATURE:
W--> 1893 <221> NAME/KEY:
     1894 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
     1895   RA patients
W--> 1896 <400> SEQUENCE: 148
     1898 tacttctgtg ccagcagtcc ccgactagcg ggagataaag gagcagtact tcgggccggg   60
E--> 1899 c   61 ← insert

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/29/2003
PATENT APPLICATION: **US/10/612,468**   TIME: 13:35:31

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

```
Seq#:1; Line(s) 21
Seq#:3; Line(s) 44
Seq#:4; Line(s) 58
Seq#:75; Line(s) 881
Seq#:77; Line(s) 910
Seq#:79; Line(s) 939
Seq#:81; Line(s) 967
Seq#:83; Line(s) 997
Seq#:85; Line(s) 1025
Seq#:87; Line(s) 1052
Seq#:89; Line(s) 1079
Seq#:91; Line(s) 1106
Seq#:93; Line(s) 1133
Seq#:95; Line(s) 1160
Seq#:97; Line(s) 1187
Seq#:99; Line(s) 1214
Seq#:101; Line(s) 1241
Seq#:103; Line(s) 1268
Seq#:105; Line(s) 1295
Seq#:107; Line(s) 1323
Seq#:109; Line(s) 1351
Seq#:111; Line(s) 1378
Seq#:113; Line(s) 1406
Seq#:115; Line(s) 1436
Seq#:117; Line(s) 1464
Seq#:119; Line(s) 1492
Seq#:121; Line(s) 1520
Seq#:123; Line(s) 1549
Seq#:125; Line(s) 1577
Seq#:127; Line(s) 1605
Seq#:129; Line(s) 1632
Seq#:131; Line(s) 1659
Seq#:133; Line(s) 1686
Seq#:135; Line(s) 1713
Seq#:137; Line(s) 1740
Seq#:139; Line(s) 1767
Seq#:141; Line(s) 1795
Seq#:143; Line(s) 1822
Seq#:145; Line(s) 1851
Seq#:147; Line(s) 1879
Seq#:149; Line(s) 1908
Seq#:151; Line(s) 1936
Seq#:153; Line(s) 1964
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/29/2003
PATENT APPLICATION: US/10/612,468   TIME: 13:35:31

Input Set : A:\EP.txt
Output Set: N:\CRF4\07292003\J612468.raw

Seq#:155; Line(s) 1992
Seq#:157; Line(s) 2020
Seq#:159; Line(s) 2047
Seq#:161; Line(s) 2075
Seq#:163; Line(s) 2103
Seq#:165; Line(s) 2131
Seq#:167; Line(s) 2159

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31
Seq#:32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,76,78,80,82,84
Seq#:86,88,90,92,94,96,98,100,102,104,106,108,110,112,114,116,118,120,122,124
Seq#:126,128,130,132,134,136,138,140,142,144,146,148,150,152,154,156,158,160
Seq#:162,164,166,168

VERIFICATION SUMMARY
PATENT APPLICATION: **US/10/612,468**

DATE: 07/29/2003
TIME: 13:35:31

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

L:7 M:283 W: Missing Blank Line separator, <120> field identifier
L:9 M:283 W: Missing Blank Line separator, <130> field identifier
L:10 M:283 W: Missing Blank Line separator, <140> field identifier
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:11 M:289 W: Identifier Missing or Out-Of-Order, <150> PRIOR APP NO
L:12 M:283 W: Missing Blank Line separator, <160> field identifier
L:18 M:283 W: Missing Blank Line separator, <220> field identifier
L:19 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:22 M:283 W: Missing Blank Line separator, <400> field identifier
L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:31 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:34 M:283 W: Missing Blank Line separator, <400> field identifier
L:42 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <400> field identifier
L:56 M:283 W: Missing Blank Line separator, <220> field identifier
L:60 M:283 W: Missing Blank Line separator, <400> field identifier
L:69 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:73 M:283 W: Missing Blank Line separator, <400> field identifier
L:82 M:283 W: Missing Blank Line separator, <220> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:109 M:283 W: Missing Blank Line separator, <220> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:137 M:283 W: Missing Blank Line separator, <220> field identifier
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
L:148 M:283 W: Missing Blank Line separator, <220> field identifier
L:151 M:283 W: Missing Blank Line separator, <400> field identifier
L:159 M:283 W: Missing Blank Line separator, <220> field identifier
L:162 M:283 W: Missing Blank Line separator, <400> field identifier
L:170 M:283 W: Missing Blank Line separator, <220> field identifier
L:173 M:283 W: Missing Blank Line separator, <400> field identifier
L:181 M:283 W: Missing Blank Line separator, <220> field identifier
L:184 M:283 W: Missing Blank Line separator, <400> field identifier
L:192 M:283 W: Missing Blank Line separator, <220> field identifier
L:195 M:283 W: Missing Blank Line separator, <400> field identifier
L:203 M:283 W: Missing Blank Line separator, <220> field identifier
L:206 M:283 W: Missing Blank Line separator, <400> field identifier
L:215 M:283 W: Missing Blank Line separator, <220> field identifier
L:218 M:283 W: Missing Blank Line separator, <400> field identifier
L:226 M:283 W: Missing Blank Line separator, <220> field identifier
L:229 M:283 W: Missing Blank Line separator, <400> field identifier
L:237 M:283 W: Missing Blank Line separator, <220> field identifier
L:240 M:283 W: Missing Blank Line separator, <400> field identifier
L:248 M:283 W: Missing Blank Line separator, <220> field identifier
L:251 M:283 W: Missing Blank Line separator, <400> field identifier
L:259 M:283 W: Missing Blank Line separator, <220> field identifier
L:262 M:283 W: Missing Blank Line separator, <400> field identifier
L:270 M:283 W: Missing Blank Line separator, <220> field identifier

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Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

```
L:273  M:283 W: Missing Blank Line separator, <400> field identifier
L:281  M:283 W: Missing Blank Line separator, <220> field identifier
L:284  M:283 W: Missing Blank Line separator, <400> field identifier
L:292  M:283 W: Missing Blank Line separator, <220> field identifier
L:295  M:283 W: Missing Blank Line separator, <400> field identifier
L:303  M:283 W: Missing Blank Line separator, <220> field identifier
L:306  M:283 W: Missing Blank Line separator, <400> field identifier
L:784  M:252 E: No. of Seq. differs, <211> LENGTH:Input:21 Found:19 SEQ:66
L:896  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:76
L:925  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:78
L:953  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:80
L:982  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:82
L:1011 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:84
L:1039 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:86
L:1066 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:88
L:1093 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:90
L:1120 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:92
L:1147 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:94
L:1174 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:96
L:1201 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:98
L:1228 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:100
L:1254 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:102
L:1282 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:104
L:1310 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:106
L:1338 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:108
L:1365 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:110
L:1392 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:112
L:1421 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:114
L:1427 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:114
L:1450 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:116
L:1456 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:116
L:1478 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:118
L:1484 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:118
L:1507 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:120
L:1534 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:122
L:1540 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:122
L:1563 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:124
L:1591 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:126
L:1619 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:128
L:1646 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:130
L:1673 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:132
L:1700 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:134
L:1727 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:136
L:1754 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:138
L:1781 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:140
L:1809 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:142
L:1837 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:144
L:1865 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:146
L:1893 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:148
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VERIFICATION SUMMARY   DATE: 07/29/2003
PATENT APPLICATION: **US/10/612,468**   TIME: 13:35:31

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

```
L:1899 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:61 SEQ:148
L:1923 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:150
L:1950 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:152
L:1978 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:154
L:2007 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:156
L:2034 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:158
L:2061 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:160
L:2089 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:162
L:2117 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:164
L:2145 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:166
L:2173 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:168
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